# Synaptotagmin-11 inhibits clathrin-mediated and bulk endocytosis

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#### **Appendix Figure S1**





A, B. Representative membrane capacitance ( $C_m$ ) traces recorded from a DRG neuron induced by a 200-ms depolarizing pulse (arrow) (A) or a 2-Hz train of twenty 100-ms depolarizing pulses (B).

C. Representative  $C_m$  trace with endocytosis overshoot recorded from a DRG neuron. Membrane conductance ( $G_m$ ), series conductance ( $G_s$ ), membrane current ( $I_m$ ), and the voltage clamp protocol  $\left(V_{m}\right)$  are shown in the lower traces.

D, E. Percentages of  $C_m$  traces with overshoot in Syt11 KD neurons (D, n = 60) *versus* control (Ctrl) (E, n = 21), scrambled shRNA (SC) (E, n = 14), and Syt11 KD neurons with Syt11 rescue (E, n = 8).

Data were collected from 4 (Sc, Rescue), 8 (Ctrl), and 16 (KD) independent experiments.

**Appendix Figure S2** 



Appendix Figure S2. Syt11 KD accelerates endocytic  $C_m$  decay and vesicle pool replenishment in DRG neurons.

A, B. Representative  $C_{\rm m}$  traces recorded from DRG neurons induced by a 200-ms  $_4$ 

depolarizing pulse (arrows). The endocytic  $C_m$  decay was fitted to a double-exponential decay function (solid blue and red, fitted curves). DRG neurons were transfected with plasmids expressing shSyt11-2 (Syt11 KD, KD) or scrambled shRNA (Sc) and  $C_m$  recording was performed 5 days after transfection. Insets show Ca<sup>2+</sup> currents recorded in the same neurons.

C, D. Time-constants of the fast and slow phases of endocytosis in Control (Ctrl), Sc, KD, and rescued (with the shSyt11-2-resistant form of Syt11) DRG neurons. Data were collected from 4 (Sc, Rescue), 8 (Ctrl), and 16 (KD) independent experiments. E. Peak  $C_{\rm m}$  changes recorded from Syt11 KD neurons, and KD neurons with Syt4 or

Syt11 rescue. Mean  $\pm$  s.e.m.; *t*-test, \*P < 0.05, \*\*P < 0.01.

F. Normalized  $C_{\rm m}$  traces recorded as in (E). Data were collected from 4 (Rescue) and 16 (KD) independent experiments.

G, H. Normalized  $\Delta C_{\rm m}$  (A) and  $C_{\rm m}$  jumps (B) induced by a 1 Hz train of 10 pulses (arrows). The normalized  $C_{\rm m}$  jumps were calculated by dividing the  $\Delta C_{\rm m}$  values in response to each pulse with that induced by the first one. Data were collected from 3 independent experiments.

All data are presented as mean  $\pm$  s.e.m. One-way ANOVA for C-F, Student's *t*-test for G and H, \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001.

### **Appendix Figure S3**





A. Fluorescence intensity of FM dye taken up into synaptic boutons. Student's *t*-test, \*\*\*P < 0.001.

B. Normalized fluorescence intensity  $(\Delta F/F_0)$  used to assess exocytosis at hippocampal synapses.

Data are presented as mean  $\pm$  s.e.m of 3 independent experiments.

#### **Appendix Figure S4**



Appendix Figure S4. Syt11 knockdown-induced fast endocytosis is partially blocked by Pitstop 1

A. Averaged  $\Delta C_m$  traces induced by 200-ms depolarization in the presence and absence of 30  $\mu$ M Pitstop 1. Endocytic inhibition by Pitstop 1 is shown by the difference in  $C_m$  traces recorded from Syt11 KD neurons (KD) and those dialyzed with Pitstop 1 (KD + Pitstop 1).

B. Endo-5s recorded as in (A).

Data are presented as mean  $\pm$  s.e.m of 3 independent experiments. Student's *t*-test, \*\*P < 0.01.



Appendix Figure S5. Syt11 expression levels in Syt11-KD and rescued DRG neurons.

A, B. Immunoblotting for Syt11 expression in control (scrambled shRNA), KD (shSyt11-2), and Syt11-rescued DRG neurons. DRG neurons were transfected with plasmids expressing scrambled shRNA, or shSyt11-2 (Syt11 KD, KD) with or without Syt11-rescue and cultured for 5 days. Transfected cells were collected with a fluorescence-activated cell-sorter and the whole-cell lysates were used for immunoblotting analysis. The results revealed similar expression levels of total Syt11 in control and rescued cells (rescued ~ 1.07-fold that in control cells).

C, D. Immunoblotting for the expression of the different Syt11 mutations in HEK293 cells. HEK293 cells were transfected with plasmids expressing full-length Syt11 (FL), or Syt11 harboring a deletion of the transmembrane domain (TMD), C2A, or C2B, or the KKAA mutant (replacing the two conserved lysine residues with alanines in the

AP-2-binding site) and immunoblotting was performed ~ 24 h after transfection. Please note that  $\Delta$ C2A was hardly detectable due to deletion of part of the antibody epitope. These experiments ruled out the possible misfolding of mutants after deletion or point mutation, and suggested that most of the rescued constructs expressed at similar levels.

Data are presented as mean  $\pm$  s.e.m of 4 independent experiments. One-way ANOVA, \*P < 0.05.